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EXAMPLE 8

D-RTR Tetramer Inhibition of N-acetyl-PGP or N-methyl-PGP
Induced PMN Polarization

The RTR complementary peptide has been shown inhibit the polarization of polymorphonuclear leukocytes activated by N-acetyl-PGP. The complementary sequence, RTR, was designed to specifically interact hydropathically with the PGP sequence in Nacetyl-PGP and, therefore, should also interact with the same sequence in N-methyl-PGP. The D-RTR tetrameric peptide N-acetyl-PGP designed inhibit to or N-methyl-PGP induced -- polymorphonuclear -- leukocyte polarization, -- but -- have -- a -- greaterstability in vivo by resisting proteolytic degradation.

A preliminary study showed that the D-RTR tetramer inhibition (n=6). The D-RTR tetramer study showed that the D-RTR tetramer $= 37\% \pm 35\%$ inhibition (n=7), 1 μ M D-RTR tetramer $= 92\% \pm 6\%$ inhibition (n=6) and 10 μ M D-RTR tetramer $= 92\% \pm 6\%$ inhibition (n=6). The D-RTR tetramer inhibited (mean \pm SD) 1 mM N-

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methyl-PGP induced polymorphonuclear leukocyte polarization as follows: 1-10 μ M D-RTR tetramer = 14% \pm 10% inhibition (n=5), 40-100 μ M D-RTR tetramer = 45% \pm 7% inhibition (n=2) and 200-800 μ M D-RTR tetramer = 100% inhibition (n=5).

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EXAMPLE 9

Results

All four complementary (antisense) peptides, containing the RTR sequence, showed substantial inhibition of N-acetyl-PGP activated polymorphonuclear leukocyte polarization (Table 1). The RTR tetrameric peptide was a powerful inhibitor of N-acetyl-PGP (ID₅₀ of 200 nM). The RTR dimer was much less potent (ID₅₀ of 105 $\mu M).$ Both monomers, RTR (ID50 of 2.5 mM) and RTRGG (ID50 of 2.1 mM), were only antagonistic at millimolar concentrations. Preincubation of the RTR tetrameric peptide with N-acetyl-PGP or neutrophils for 5 min did not change the results described above. antisense peptide, ASA tetramer, failed to show any additional inhibition of polymorphonuclear leukocytes activated by N-acetyl-PGP.

TABLE I

Complementary Peptide Inhibition of N-acetyl-PGP Activated PMN

Polarization

Complementary	Antagonist	p-value
Peptides	Activity (ID ₅₀)	
RTR tetramer	200 nM ± 75 nM	<0.001
RTR dimer	105 μM ± 68 μM	0.001
RTR monomer	2.5 mM ± 1.2 mM	<0.001
RTRGG monomer	2.1 mM ± 0.8 mM	<0.001
ASA tetramer	None, ≤ 4 mM	

5 * Untreated PMNs (negative control) produced a polarization

response of 7.8% ± 4.4% (n = 41). PMNs activated with 500 µM N-acetyl-PGP (positive control) produced a polarization response of 56.5% ± 16.4% (n = 41). This chemoattractant concentration was selected from the linear portion of the dose response curve, yielding approximately 50% polarization after subtraction of the negative control values. Antagonistic activity (ID50, mean ± standard deviation) was interpolated from five dose response curves for each complementary peptide.